HCV Genome and Recombinant Proteins

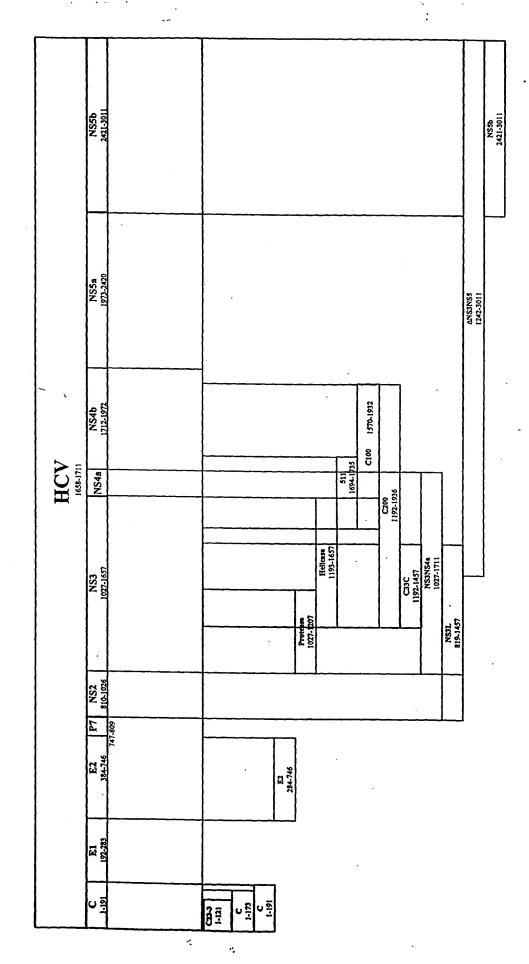


FIG. 1

										A GCG				10 Q CAG
										S AGC				
										I ATT				
										G GGG				
										A GCG				
										Q CAA				
										T ACA				
										H CAC				
										S AGC				
										G GGG				L TTG
														160 C TGC
										I ATC				
L	E	T	т	180 M	R	s	P	v	F	T	D	N	s	190

2	^	$\sim$

P P V V P Q S F Q V A H L H A CCA CCA GTA GTG CCC CAG AGC TTC CAG GTG GCT CAC CTC CAT GCT

210 220

P T G S G K S T K V P A A Y A CCC ACA GGC AGC GGC AAA AGC ACC AAG GTC CCG GCT GCA TAT GCA

230

A Q G Y K V L V L N P S V A A GCT CAG GGC TAT AAG GTG CTA GTA CTC AAC CCC TCT GTT GCT GCA

240 250

T L G F G A Y M S K A H G I D ACA CTG GGC TTT GGT GCT TAC ATG TCC AAG GCT CAT GGG ATC GAT

260

P N I R T G V R T I T T G S P CCT AAC ATC AGG ACC GGG GTG AGA ACA ATT ACC ACT GGC AGC CCC

270 280

I T Y S T Y G K F L A D G G C ATC ACG TAC TCC ACC TAC GGC AAG TTC CTT GCC GAC GGC GGG TGC

290

S G G A Y D I I I C D E C H S TCG GGG GGC GCT TAT GAC ATA ATA ATT TGT GAC GAG TGC CAC TCC

300 310

T D A T S I L G I G T V L D Q ACG GAT GCC ACA TCC ATC TTG GGC ATT GGC ACT GTC CTT GAC CAA

320

A E T A G A R L V V L A T A T GCA GAG ACT GCG GGG GCG AGA CTG GTT GTG CTC GCC ACC GCC ACC

330 340

P P G S V T V P H P N I E E V CCT CCG GGC TCC GTC ACT GTG CCC CAT CCC AAC ATC GAG GAG GTT

350

A L S T T G E I P F Y G K A I GCT CTG TCC ACC GGA GAG ATC CCT TTT TAC GGC AAG GCT ATC

360 370

P L E V I K G G R H L I F C H
CCC CTC GAA GTA ATC AAG GGG GGG AGA CAT CTC ATC TGT CAT

380

S K K K C D E L A A K L V A L TCA AAG AAG AGG TGC GAC GAA CTC GCC GCA AAG CTG GTC GCA TTG

				390										400
G	I	N	A	v	A	Y	Y	R	G	L	D	V	S	v
GGC	ATC	AAT	GCC	GTG	GCC	TAC	TAC	CGC	GGT	CTT	GAC	GTG	TCC	GTC
									410					
												D		L
ATC	CCG	CCC	ATC	GGC	GAT	GTT	GTC	GTC	GTG	GCA	ACC	GAT	GCC	CTC
				420				_	_		_	_	~	430
M	T	G	Y	T	G	D	F	D	S	V	1	D	maa.	N
ATG	ACC	GGC	TAT	ACC	GGC	GAC	TTC	GAC	TCG	GTG	ATA	GAC	IGC	AAI
			•	•					440					
m	~	7.7	·	^	T	77	Ъ	E.		Ť.	מ	P	т	F
J. C.C.	. С	CTC	y C.C.	עאַכּ	ארש	GTC	CAT	ים ביים די	AGC	CTT	GAC	CCT	ACC	TTC
ACG	161	GIC	ACC	CAG	ACA	GIC	OAI	110	7100	<b></b>	00	-		
				450										460
т	I	E	т		T	L	P	. Q	D	A	V	s	R	T
ACC	ATT	GAG	ACA	ATC	ACG	CTC	CCC	CAA	GAT	GCT	GTC	TCC	CGC	ACT
,														
									470					
Q	R	R	G	R	T	G	R	G	K	P	G	I	Y	R
CAA	CGT	CGG	GGC	AGG	ACT	GGC	AGG	GGG	AAG	CCA	GGC	ATC	TAC	AGA
			•											400
				480		_	_	_	_		_	-		490
F	V	A	P	G	E	R	D C C C	S	G	N MC	r mmc	D	TICC	TCC
TTT	GTG	GCA	CCG	فافافا	GAG	CGC	CCC	TCC	GGC	AIG	110	GAC	100	100
									500					
37	т.	C	<b>R</b> .	C	v	ח	Δ	G			W	Y	E	L
GTC	רער טער	TGT	GAG	тас	TAT	GAC	GCA	GGC	TGT	GCT	TGG	TAT	GAG	CTC
010	CIC	101	00	100			00							
				510										520
T	P	A	E	T	T	v	R	L	R	A	Y	M	N	${f T}$
ACG	CCC	GCC	GAG	ACT	ACA	GTT	AGG	CTA	CGA	GCG	TAC	ATG	AAC	ACC
									530		•			_
	G											W		
CCG	GGG	CTT	, CCC	GTG	TGC	CAG	GAC	CAT	CTI	GAA	TTT	TGG	GAG	GGC
														EEA
		_		540			_	_		••	-		c	550
V	F	T	G	L	T	H		ט	A	H	r anana	, Cany	שכר שכר	Q
GTC	TTT	ACA	GGC	CTC	: ACI	CAT	ATA	GAT	GCC	CAC	, 111	CIA	. 100	CAG
									560	,				
T.	v	^	c	a	127	M	۲.	D			v	Δ	Y	Q
ב ה	אמ. ממ.	י קאר ע	ים דים א	י נכננת י נכננת	CDC E	ן. אסע	Cului Ti	CCT	<u>.</u> ' ፐልር	CTY	GTA	GCG	TAC	CAA
ACF	· 'ww	, cac	, AG1	. 330	, one									
				570	)									580
A	T	v	С			A	Q	A	P	P	P	S	W	D
GC	ACC	GTO	TGC	GCI	' AGG	GCI	CAA	GCC	: cc	r ccc	CCA	TCG	TGG	GAC

Q M W K C L I R L K P T L H G CAG ATG TGG AAG TGT TTG ATT CGC CTC AAG CCC ACC CTC CAT GGG

600

P T P L L Y R L G A V Q N E I CCA ACA CCC CTG CTA TAC AGA CTG GGC GCT GTT CAG AAT GAA ATC

620

TLTHPVTKYIMTCMS ACC CTG ACG CAC CCA GTC ACC AAA TAC ATC ATG ACA TGC ATG TCG

> 640 630

ADLEVVTSTWVLVGG GCC GAC CTG GAG GTC GTC ACG AGC ACC TGG GTG CTC GTT GGC GGC

650

V L A A L A A Y C L S GTC CTG GCT GCT TTG GCC GCG TAT TGC CTG TCA ACA GGC TGC GTG

660

IVGRVVLSGKPAII GTC ATA GTG GGC AGG GTC GTC TTG TCC GGG AAG CCG GCA ATC ATA

680

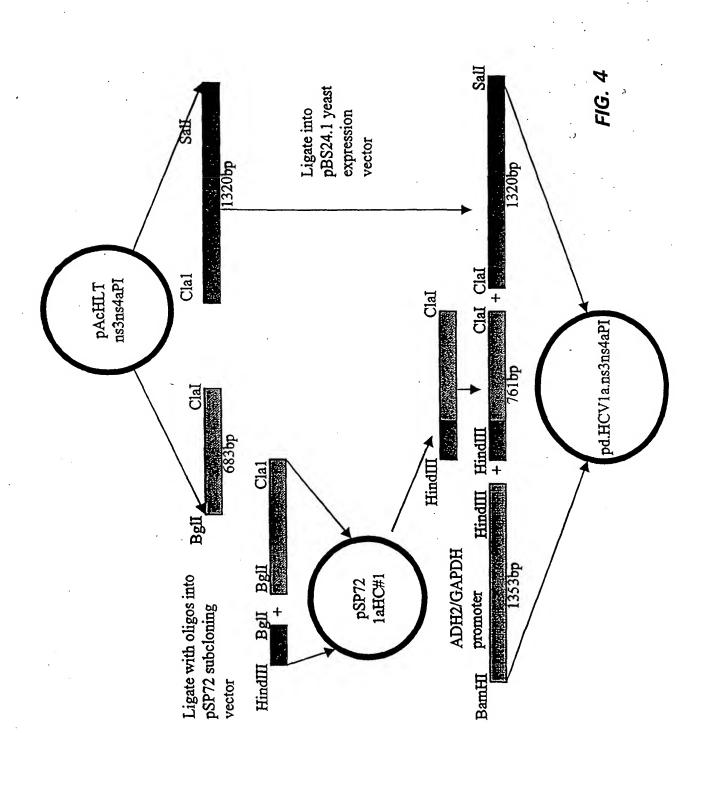
E V L Y R E F D E CCT GAC AGG GAA GTC CTC TAC CGA GAG TTC GAT GAG ATG GAA GAG

686

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. TGC

FIG. 3D



## **MEFA 12 Antigen Construct**

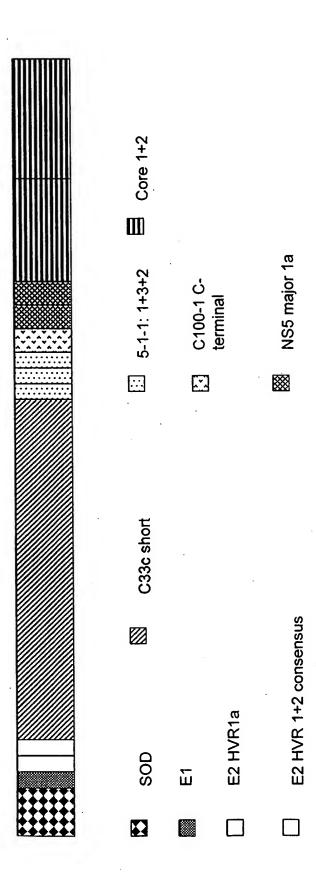


FIG. 5

1									10						
M ATG					V GTT									V GTT	45
					F TTC									30 V GTG	90
					I ATT										135
F	Н	V	Н	50 E	F	G	D	N	Т	A	G	С	Т	60 S	180
					N AAT										225
					H CAT										270
					R AGA										315
					Q CAA									120 A GCA	360
A GCC					S TCT									G GGT	405

		_	 140	_	_	_	_	_	_	_	 ~	150	
									T ACG			S TCT	450
									A GCT			A GCT	495
									P CCG			180 A GCA	540
A GCT	_								P CCC			A GCA	585
									A GCT			210 D GAT	630
P CCT									T ACC			P CCC	675
									A GCC			240 C· TGC	720
									D GAC			S TCC	765
												270 Q CAA	
									L CTC			T ACC	855
									N AAC			300 V GTT	900

		T ACC						I ATC	945
		320 I ATC							990
		C TGC							1035
		350 V GTG							1080
		G GGC						L CTC	1125
		380 T ACC							1170
		S TCC							1215
		410 E GAG							1260
		E GAG							1305
		440 G GGC				P CCG	I ATC		1350
		V GTG				E GAG		E GAG	1395

C TGC	S TCA	Q CAA	A GCT	470 A GCC	P CCA	Y TAT	I ATC	E GAA	Q CAA		V GTA		480 A GCT	1440
H CAC			K AAG								N AAT		Q CAA	1485
		V GTG							L TTA			F TTT	510 D GAT	1530
E GAG				C TGC					520 A GCC		E GAG		G GGG	1575
			A GCG								G GGC		540 L CTC	1620
G GGG		L CTG	R CGC	R CGG					550 G GGC			V GTG	Q CAG	1665
W TGG			R CGG						S TCC	G GGG			570 V GTT	1710
S TCC			H CAC	Y TAC					580 S TCC				Q CAG	1755
			V GTT										600 V GTG	1800
			K AAA											1845
			R CGG										630 R CGG	1890

D Y N P P L V E T W K K P D CCG GAC TAT AAC CCC CCG CTA GTG GAG ACG TGG AAA AAG CCC GAC 1935 660 650 V H G R K T K N ₽ V E Ρ TAC GAA CCA CCT GTG GTC CAT GGC AGA AAG ACC AAA CGT AAC ACC 670 P Q D V K P G G G Q I F R AAC CGG CGG CCG CAG GAC GTC AAG TTC CCG GGT GGC GGT CAG ATC 690 680 V Y L L P R R G P R L G GTT GGT GGA GTT TAC TTG TTG CCG CGC AGG GGC CCT AGA TTG GGT 700 A T R K T S P I P K A R R GTG CTC GCG ACG AGA AAG ACT TCC CCT ATC CCC AAG GCT CGT CGG 720 710 WAQPGYPWPL R  ${f T}$ CCC GAG GGC AGG ACC TGG GCT CAG CCC GGT TAC CCT TGG CCC CTC 730 T G K S W G K Y G N K D R R S TAT GGC AAT AAG GAC AGA CGG TCT ACA GGT AAG TCC TGG GGT AAG 740 N R PRKTKRN Y P W $\mathbf{T}$ CCA GGG TAC CCT TGG CCA AGA AAG ACC AAA CGT AAC ACC AAC CGG 2250 760 P Q D V K F P G G G I V 0 CGG CCG CAG GAC GTC AAG TTC CCG GGT GGC GGT CAG ATC GTT GGT 770 780 G P R L G V L V Y L L P R R GGA GTT TAC TTG TTG CCG CGC AGG GGC CCT AGA TTG GGT GTG CTC 2340 790 PKARRPE K T S P I Α R GCG ACG AGA AAG ACT TCC CCT ATC CCC AAG GCT CGT CGG CCC GAG 2385

820 -

N K D R R S T G K S W G K P G AAT AAG GAC AGA CGG TCT ACA GGT AAG TCC TGG GGT AAG CCA GGG 2475

829

Y P W P OC TAC CCT TGG CCC TAA TGAGTCGAC

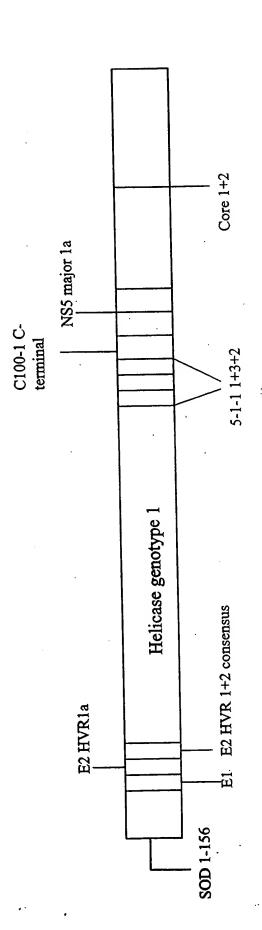


FIG. 7

A T K A V C V L K G D G P V ATG GCT ACA AAG GCT GTT TGT GTT TTG AAG GGT GAC GGC CCA GTT 20 30 FEQKESNGPV QGIIN CAA GGT ATT ATT AAC TTC GAG CAG AAG GAA AGT AAT GGA CCA GTG 40 WGSIKGL T E AAG GTG TGG GGA AGC ATT AAA GGA CTG ACT GAA GGC CTG CAT GGA 50 F G D N T A G C T V H E TTC CAT GTT CAT GAG TTT GGA GAT AAT ACA GCA GGC TGT ACC AGT 70 PHFNPLS R K GCA GGT CCT CAC TTT AAT CCT CTA TCC AGA AAA CAC GGT GGG CCA H V G D L G N V T A E R AAG GAT GAA GAG AGG CAT GTT GGA GAC TTG GGC AAT GTG ACT GCT 100 D K D G V A D V S I E GAC AAA GAT GGT GTG GCC GAT GTG TCT ATT GAA GAT TCT GTG ATC 110 G D H C I I G R T L V V TCA CTC TCA GGA GAC CAT TGC ATC ATT GGC CGC ACA CTG GTG GTC 130 E K A D D L G K G G N E E CAT GAA AAA GCA GAT GAC TTG GGC AAA GGT GGA AAT GAA GAA AGT 140 150 T: K T G N A G S R L A C G V I ACA AAG ACA GGA AAC GCT GGA AGT CGT TTG GCT TGT GGT GTA ATT 160 I A Q N L N S G C N C S I Y GGG ATC GCC CAG AAT TTG AAT TCT GGT TGC AAT TGC TCT ATC TAT 170 180 ITGHRMAWKLG CCC GGC CAT ATA ACG GGT CAC CGC ATG GCA TGG AAG CTT GGT TCC 190 A R T T S G F V S L F A P GCC GCC AGA ACT ACC TCG GGC TTT GTC TCC TTG TTC GCC CCA GGT

	N AAC	E	T	н		${f T}$	G	G			
	G GGG									S AGC	
	L TTG										
	R CGA										
	Q CAG										
	K AAA										
	V GTG										
	A										
	G G										330 T ACG
	Y TAC										
	D GAC										
	I ATC							L			
	A G GCG										
	T C ACT							E			
		440		3	,	•	. •				420

	T ACC													
	V GTA												S TCA	
	K AAG													
	A GCC													
	S AGC													480 T ACC
	Y TAT													
	T ACC													
	T ACA													R CGT
	G GGC													
	P													
C TGT	E GAG					G GGC		A GCT				L CTC		570 P CCC
	E GAG													G GGG
	p CCC				D									600 F TTT
	G G G									L				K AAG
		~	. 15	620			v		17	*	v	^		630 T

CAG AGT	GGG	GAG	AAC	CTT	CCT	TAC	CTG	GTA	GCG	TAC	CAA	GCC	ACC
V C													
W K													
P L CCC CTG													
T H ACG CAC													
L E													
P D CCT GAC													
C S TGC TCT													
E Q GAG CAG		K											
P A													
E M GAG ATG				S									780 A GCT
Q V CAG GTA													
D N GAT AAT													810 Y TAT
E A GAG GCC													
I E ATT GAG													840 I ATA

P Q D V K F P G G Q I

AAC CGA CGG CCG CAG GAC GTC AAG TTC CCG GGT GGC GGT CAG ATC

V G R R G P P I P K A R R P E . GTT GGT CGC AGG GGC CCT CCT ATC CCC AAG GCT CGT CGG CCC GAG

1070

G R T W A Q P G Y P W P L Y G GGC AGG ACC TGG GCT CAG CCC GGT TAC CCT TGG CCC CTC TAT GGC

1090

N K D R R S T G K S W G K P G AAT AAG GAC AGA CGG TCT ACC GGT AAG TCC TGG GGT AAG CCA GGG

1099

Y P W P TAT CCT TGG CCC

FIG. 8F

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hSOD- (1-154)	CORE	CORE	c33c	5-1-1 type 1	5-1-1 type 3	5-1-1 type 2	C-100	Q-100	NS5	NSS
AMINO 10	0	10	1192	1694	1694	1694	1901	1901	2278	2278
ACIDS	53	53	1457	1735	1735	1735	1940	1940	2310	2310

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## MEFA-5 ANTIGEN

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hSOD- (1-154)	CORE	CORE	E1	ដ	යි3ී	5-1-1 type 1	5-1-1 type 3	5-1-1 type 2	C-100	NSS	NSS
AMINO	10 -	10 -	303 -	405 -	1192 -	1689 -	1689 -	1689 -	1901 -	2278 -	2278 -
ACIDS	53	23	320	444	1457	1735	1735	1735	1940	2313	2313

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## MEFA-6 ANTIGEN

hSOD- (1-154)	Ω	<b>E2</b>	ය3ිය	5-1-1 type 1	5-1-1 type 3	5-1-1 type 2	C-100	NS5	NS5	CORE	CORE
AMINO	303 -	405 -	1192 -	1689 -	1689 -	1689 -	1901 -	2278-	2278-	10-	10-
ဗ	320	444	1457	1735	1735	1735	1940	2313	2313	33	23